



Sequence Listing

<110> Hirokazu Matsumoto
Jiro Noguchi
Mioko Harada
Masaaki Mori

<120> Body weight gain inhibitor

<130> 61536 (46342)

<140> 10/500,175

<141> 2004-06-25

<150> PCT/JP01/13781

<151> 2002-12-27

<150> JP2001-403260

<151> 2001-12-28

<150> JP2002-93096

<151> 2002-03-28

<160> 150

<210> 1

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 1

atcgattaca atgcaggccg ctgggcaccc ag 32

<210> 2

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 2

actagtgcc ttcagcaccc caatatgctg cg 32

<210> 3

<211> 1023

<212> DNA

<213> Homo sapiens

<400> 3

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ctccgagcca	ctgccgttcc	tctatgtgct	cctgcccgcc	gtgtactccg	ggatctgtgc	180
tgtggggctg	actggcaaca	cggccgtcat	ccttgtaatc	ctaagggcgc	ccaagatgaa	240
gacggtgacc	aacgtgttca	tctgaacct	ggccgtcgcc	gacgggctct	tcacgctggg	300

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actgcccgtc aacatcgcgg agcacctgct gcagtactgg cccttcgggg agctgctctg 360
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<210> 4

<211> 333

<212> PRT

<213> Homo sapiens

<400> 4

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20          25          30
His Asn Ala Thr Phe Ser Glu Pro Leu Pro Phe Leu Tyr Val Leu Leu
35          40          45
Pro Ala Val Tyr Ser Gly Ile Cys Ala Val Gly Leu Thr Gly Asn Thr
50          55          60
Ala Val Ile Leu Val Ile Leu Arg Ala Pro Lys Met Lys Thr Val Thr
65          70          75          80
Asn Val Phe Ile Leu Asn Leu Ala Val Ala Asp Gly Leu Phe Thr Leu
85          90          95
Val Leu Pro Val Asn Ile Ala Glu His Leu Leu Gln Tyr Trp Pro Phe
100         105         110
Gly Glu Leu Leu Cys Lys Leu Val Leu Ala Val Asp His Tyr Asn Ile
115         120         125
Phe Ser Ser Ile Tyr Phe Leu Ala Val Met Ser Val Asp Arg Tyr Leu
130         135         140
Val Val Leu Ala Thr Val Arg Ser Arg His Met Pro Trp Arg Thr Tyr
145         150         155         160
Arg Gly Ala Lys Val Ala Ser Leu Cys Val Trp Leu Gly Val Thr Val
165         170         175
Leu Val Leu Pro Phe Phe Ser Phe Ala Gly Val Tyr Ser Asn Glu Leu
180         185         190
Gln Val Pro Ser Cys Gly Leu Ser Phe Pro Trp Pro Glu Gln Val Trp
195         200         205
Phe Lys Ala Ser Arg Val Tyr Thr Leu Val Leu Gly Phe Val Leu Pro
210         215         220
Val Cys Thr Ile Cys Val Leu Tyr Thr Asp Leu Leu Arg Arg Leu Arg
225         230         235         240
Ala Val Arg Leu Arg Ser Gly Ala Lys Ala Leu Gly Lys Ala Arg Arg
245         250         255
Lys Val Thr Val Leu Val Leu Val Val Leu Ala Val Cys Leu Leu Cys
260         265         270
Trp Thr Pro Phe His Leu Ala Ser Val Val Ala Leu Thr Thr Asp Leu
275         280         285
Pro Gln Thr Pro Leu Val Ile Ser Met Ser Tyr Val Ile Thr Ser Leu
290         295         300

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Ser Tyr Ala Asn Ser Cys Leu Asn Pro Phe Leu Tyr Ala Phe Leu Asp
 305 310 315 320
 Asp Asn Phe Arg Lys Asn Phe Arg Ser Ile Leu Arg Cys
 325 330

<210> 5
 <211> 687
 <212> RNA
 <213> Artificial Sequence

<220>
 <223> Riboprobe

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 ucaggcacga guuggcguag cugaggcugg ugaugacgua ggacauacug augaccagug 180
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 cuagagccuu ggcuccagag cggagccgca cggcccgcag ccugcgcagg aggucugugu 360
 agagcacaca gauggugcac acgggcagca cgaagcccag gaccaacgug uagacacggc 420
 uggccuugaa ccagaccugc ucgggccacg ggaagcucag cccacagcuu gggaccugca 480
 gcucguugcu guagacgcca gcgaaagaga agaagggcag aaccaggacc gugacgcca 540
 gccagacaca caggcuggcg accuucgccc cccgguaggu gcgccagggc auguggcggg 600
 accucacggg ggccagcacc accagguauc gguccacgcu caucacggcu aggaaguaga 660
 ugcuggagaa gauguuguag uggucga 687

<210> 6
 <211> 17
 <212> PRT
 <213> Sus scrofa

<400> 6
 Trp Tyr Lys His Thr Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala

<210> 7
 <211> 438
 <212> DNA
 <213> Homo sapiens

<220>
 <221> misc_feature
 <222> 408
 <223> n is a, c, g, or t

<400> 7
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 tgtaccacgc gccggagggc agcggcagca ggagcagaag cagcagcagt gccagccgcg 120
 gccggctcgc gggagcccc cgcctcccctg ggcgccacgc cagggcgctc gcgtcgacgg 180
 ccgcccggcg gggcgggcca cgaaccggct cggctgggggt tgggcgcgca gtggagttag 240
 gacgccacag taccggagcg caggaggctg gaggcgagcc gtgggtcccc tgcaggccca 300
 gctataaccg ctcggtggcc ccgcctcgtt ccgccccctc agtaccgctg ggctccccag 360
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 ttgagcacct cactgagt 438

<210> 8

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<211> 264
<212> DNA
<213> Homo sapiens

<400> 8
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gcagcagtgc cagccgcggc cggtcgcggg gagccccccg ctcccctggg cgccacgcca 180
gggcgctcgc gtcgacggcc gcccggcggg gcggggccacg aaccggctcg gctgggtttg 240
ggcgcgagc ggagttggga cgcc 264

<210> 9
<211> 424
<212> DNA
<213> Homo sapiens

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gcagcagtgc cagccgcggc cggtcgcggg gagccccccg ctcccctggg cgccacgcca 180
gggcgctcgc gtcgacggcc gcccggcggg gcggggccacg aaccggctcg gctgggtttg 240
ggcgcgagc ggagttggga cgcccaggta ccggagcgca ggaggctgga ggcgagccgt 300
gggtcccctg caggcccagc tataaccgct cgggtggcccc gcctcgttcc gccccctcag 360
taccgctggg ctcccagat ggggggaggg acggagggag gagagggaac cctggcagct 420
ggcg 424

<210> 10
<211> 375
<212> DNA
<213> Homo sapiens

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tcccctgggc gccacgcagg gctacagcgt cgacggccgc ccgcggggcc atcgcaaccg 180
gctcggtcgg gtttgggcgc gcagtggagt tgggacgccc aggtaccgga gcgcaggagg 240
ctggaggcga gccgtgggtc cctcgcaggc ccagctataa ccgctcgggtg gccccgcctc 300
gttccgcccc ctcagtaccg ctgggctccc cagaatgggg gagggacgga gggaggagag 360
ggaaccctgg cagct 375

<210> 11
<211> 260
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<222> 2, 61, 147, 189, 213, 237, 249
<223> n is a, c, g, or t

<400> 11
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cccctgctac gcttactgct gcttctnctc ttgctacctc tgcccgccag cgccctggtag 180
aagcacgtng cgagccctcg ctatcacaca gtnggtcgtg cctccgggct gctcatnggg 240
ctgcgccgnt cgtcctacct 260

<210> 12

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<211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 12
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<210> 13
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 13
 tctccacag ctctgaacc cacg 24

<210> 14
 <211> 375
 <212> DNA
 <213> Homo sapiens

<400> 14
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 gacaccctct cccccgaacc cgcagcccgc gaggctcctc tcctgctgcc ctcggtgggtt 360
 caggagctgt gggag 375

<210> 15
 <211> 125
 <212> PRT
 <213> Homo sapiens

<400> 15
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 Arg Gly Ala Pro Ala Ser Arg Pro Arg Leu Ala Leu Leu Leu Leu
 35 40 45
 Leu Leu Leu Pro Leu Pro Ser Gly Ala Trp Tyr Lys His Val Ala Ser
 50 55 60
 Pro Arg Tyr His Thr Val Gly Arg Ala Ala Gly Leu Leu Met Gly Leu
 65 70 75 80
 Arg Arg Ser Pro Tyr Leu Trp Arg Arg Ala Leu Arg Ala Ala Gly
 85 90 95
 Pro Leu Ala Arg Asp Thr Leu Ser Pro Glu Pro Ala Ala Arg Glu Ala
 100 105 110
 Pro Leu Leu Leu Pro Ser Trp Val Gln Glu Leu Trp Glu
 115 120 125

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<210> 16
<211> 23
<212> PRT
<213> Homo sapiens

<400> 16
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Ala Gly Leu Leu Met Gly Leu
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<210> 17
<211> 30
<212> PRT
<213> Homo sapiens

<400> 17
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1 5 10 15
Ala Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp
20 25 30

<210> 18
<211> 69
<212> DNA
<213> Homo sapiens

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atggggctg 69

<210> 19
<211> 90
<212> DNA
<213> Homo sapiens

<400> 19
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atggggctgc gtcgctcacc ctatctgtgg 90

<210> 20
<211> 29
<212> PRT
<213> Homo sapiens

<400> 20
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1 5 10 15
Ala Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu
20 25

<210> 21
<211> 28
<212> PRT
<213> Homo sapiens

<400> 21
Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala

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						Pro
						Tyr

<210> 22
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 <212> PRT
 <213> Homo sapiens

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				20				25						

<210> 23
 <211> 26
 <212> PRT
 <213> Homo sapiens

<400> 23														
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				20				25						

<210> 24
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 <212> PRT
 <213> Homo sapiens

<400> 24														
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				20				25						

<210> 25
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 <212> PRT
 <213> Homo sapiens

<400> 25														
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				20										

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 <211> 87
 <212> DNA
 <213> Homo sapiens

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<210> 27

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<211> 84
<212> DNA
<213> Homo sapiens

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atggggctgc gtcgctcacc ctat 84

<210> 28
<211> 81
<212> DNA
<213> Homo sapiens

<400> 28
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atggggctgc gtcgctcacc c 81

<210> 29
<211> 78
<212> DNA
<213> Homo sapiens

<400> 29
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atggggctgc gtcgctca 78

<210> 30
<211> 75
<212> DNA
<213> Homo sapiens

<400> 30
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atggggctgc gtcgc 75

<210> 31
<211> 72
<212> DNA
<213> Homo sapiens

<400> 31
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atggggctgc gt 72

<210> 32
<211> 999
<212> DNA
<213> Homo sapiens

<400> 32
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ctggccgtcg accactacaa catctttctc agcatctact tcctagccgt gatgagcgtg 420
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cggggggcca aggtcgccag cctgtgtgtc tggctgggcg tcacggtcct gggtctgccc 540
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ttcccgtggc ccgagcgggt ctggttcaag gccagccgtg tctacacttt gggtcctggc 660
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<210> 33

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 33

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<210> 34

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 34

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<210> 35

<211> 1102

<212> DNA

<213> Homo sapiens

<400> 35

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<210> 36
 <211> 24
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<220>
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<400> 36
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<210> 37
 <211> 24
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<220>
 <223> Primer

<400> 37
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<210> 38
 <211> 609
 <212> DNA
 <213> Homo sapiens

<400> 38
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 gaattctag 609

<210> 39
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 39
 agcggtagtg agggggcgga acga 24

<210> 40
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

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<400> 40
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<210> 41
<211> 719
<212> DNA
<213> Homo sapiens

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gttcgtggcc cgtcccgccg ggcggccgtc gacgcgagcg ccctggcgtg gcgcccaggg 180
gagcgggggg ctcccgcgag ccggccgcgg ctggcactgc tgctgcttct gctcctgctg 240
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ccggagtccc tggacttcag cggagctggc cagagacttc ggagagacgt ctcccgccca 600
gcggtggacc ccgcagcaaa ccgccttggc ctgccctgcc tggcccccg accgttctga 660
cagcgtcccc cgcccgcccg tggcgctcc gcgcctgacc caggaggagt ggccgcgcg 719

<210> 42
<211> 165
<212> PRT
<213> Homo sapiens

<400> 42
Leu Ala Trp Arg Pro Gly Glu Arg Gly Ala Pro Ala Ser Arg Pro Arg
1 5 10 15
Leu Ala Leu Leu Leu Leu Leu Leu Leu Pro Leu Pro Ser Gly Ala
20 25 30
Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
35 40 45
Ala Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp Arg Arg
50 55 60
Ala Leu Arg Ala Ala Ala Gly Pro Leu Ala Arg Asp Thr Leu Ser Pro
65 70 75 80
Glu Pro Ala Ala Arg Glu Ala Pro Leu Leu Leu Pro Ser Trp Val Gln
85 90 95
Glu Leu Trp Glu Thr Arg Arg Arg Ser Ser Gln Ala Gly Ile Pro Val
100 105 110
Arg Ala Pro Arg Ser Pro Arg Ala Pro Glu Pro Ala Leu Glu Pro Glu
115 120 125
Ser Leu Asp Phe Ser Gly Ala Gly Gln Arg Leu Arg Arg Asp Val Ser
130 135 140
Arg Pro Ala Val Asp Pro Ala Ala Asn Arg Leu Gly Leu Pro Cys Leu
145 150 155 160
Ala Pro Gly Pro Phe
165

<210> 43
<211> 24
<212> DNA
<213> Artificial Sequence

<220>

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<223> Primer

<400> 43

acagataggg tgagcgacgc agcc 24

<210> 44

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 44

tgagcgacgc agcccatga gcag 24

<210> 45

<211> 235

<212> DNA

<213> Sus scrofa

<400> 45

cgacacccct	gcgcccagac	cctccggagc	cagttcctgg	tccgccccgc	cgggagccgt	60
cagcatgaac	ccccgggcac	gcggcatggg	agcgcggggc	ccgggaccgg	gggccactgc	120
gaggcgccgg	ctgctggcat	tgctgttact	gctgctgctg	ctgccgctgc	ccgcccgtgc	180
ctggtacaag	cacacggcga	gtccccgcta	ccacacggtg	ggccgcgccc	cgggc	235

<210> 46

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 46

cagcggcagc agcagcagca gtaa 24

<210> 47

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 47

cagcagtaac agcaatgccca gcag 24

<210> 48

<211> 156

<212> DNA

<213> Sus scrofa

<400> 48

ctgtagcctc	ccgcgctgcg	gcttccccgac	acccctgcgc	ccagaccctc	cggagccagt	60
tcctggtccg	ccccgccggg	agccgtcagc	atgaaccccc	gggcacgcgg	catgggagcg	120
cggggcccgg	gaccgggggc	cactgcgagg	cgccgg			156

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<210> 49
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 49
 cggctgctgg cattgctggt actg 24

<210> 50
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 50
 cgcccgtgcc tgggtacaagc aca 23

<210> 51
 <211> 588
 <212> DNA
 <213> Sus scrofa

<400> 51
 cggcgagtcc ccgctaccac acggtggggc gcgcgcgcggg cctgctcatg gggctgcgcc 60
 gctcgcccta catgtggcgc cgcgcgctgc gcccggcggc cgggcccctg gcctgggaca 120
 ctttcggcca ggacgtgccc cctcggggac cctccgccag gaacgccctc tctccggggc 180
 ccgcccctcg cgacgtcccg ctgcttcccc ccgggggtta gacactgtgg caggtgcgac 240
 gcggaagctt ccgctccggg atcccggta gtgcgccccg cagcccgcgc gcccgggggt 300
 ccgagccgca accggaattg ggcgcctctt cctggacctc ggcgagtag accagagcct 360
 tcggagagtc ttcagctcag cgggtggtctg cgcagggaac cgcttcgcc agccccgcc 420
 tcgccccagc gtcagagccg acctgatcgc ggccccggcg gcgcggcccc gcgcctggcc 480
 cccgcggagt ctcttcgcgc ccccaggccg gccgtctggt caataaaacc cgcctagtgc 540
 ctgcgaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaaaaaaa 588

<210> 52
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 52
 ttcccagacac ccctgcgccc agac 24

<210> 53
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

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<400> 53
 gggctggcga aggcggttcc ctgc 24

<210> 54
 <211> 565
 <212> DNA
 <213> Sus scrofa

<400> 54
 cctccggagc cagttcctgg tccgccccgc cgggagccgt cagcatgaac ccccgggcac 60
 gcggcatggg agcgcggggc ccgggaccgg gggccactgc gaggcgccgg ctgctggcat 120
 tgctgttact gctgctgctg ctgccgctgc ccgcccgtgc ctggtacaag cacacggcga 180
 gtccccgcta ccacacggtg ggccgcgcgc cgggcctgct catggggctg cgccgctcgc 240
 cctacatgtg gcgcgcgcgc ctgcgcgccg cggccgggccc cctggcctgg gacactttcg 300
 gccaggacgt gccccctcgg ggaccctccg ccaggaacgc cctctctccg gggcccggccc 360
 ctgcgcagcgc tccgctgctt cccccggggg ttcagacact gtggcagggtg cgacgcggaa 420
 gcttccgctc cgggatcccc gtcagtgcgc cccgcagccc gcgcgcccg ggggtccgagc 480
 cgcaaccgga attgggcgcc tcttcctgga cctcggcgga gtagaccaga gccttcggag 540
 agtcttcagc tcagcgggtg tctgc 565

<210> 55
 <211> 159
 <212> PRT
 <213> Sus scrofa

<400> 55
 Met Asn Pro Arg Ala Arg Gly Met Gly Ala Arg Gly Pro Gly Pro Gly
 1 5 10 15
 Ala Thr Ala Arg Arg Arg Leu Leu Ala Leu Leu Leu Leu Leu Leu Leu
 20 25 30
 Leu Pro Leu Pro Ala Arg Ala Trp Tyr Lys His Thr Ala Ser Pro Arg
 35 40 45
 Tyr His Thr Val Gly Arg Ala Gly Leu Leu Met Gly Leu Arg Arg
 50 55 60
 Ser Pro Tyr Met Trp Arg Arg Ala Leu Arg Pro Ala Ala Gly Pro Leu
 65 70 75 80
 Ala Trp Asp Thr Phe Gly Gln Asp Val Pro Pro Arg Gly Pro Ser Ala
 85 90 95
 Arg Asn Ala Leu Ser Pro Gly Pro Ala Pro Arg Asp Ala Pro Leu Leu
 100 105 110
 Pro Pro Gly Val Gln Thr Leu Trp Gln Val Arg Arg Gly Ser Phe Arg
 115 120 125
 Ser Gly Ile Pro Val Ser Ala Pro Arg Ser Pro Arg Ala Arg Gly Ser
 130 135 140
 Glu Pro Gln Pro Glu Leu Gly Ala Ser Ser Trp Thr Ser Ala Glu
 145 150 155

<210> 56
 <211> 23
 <212> PRT
 <213> Sus scrofa

<400> 56
 Trp Tyr Lys His Thr Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala Gly Leu Leu Met Gly Leu
 20

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<210> 57
 <211> 30
 <212> PRT
 <213> Sus scrofa

<400> 57
 Trp Tyr Lys His Thr Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Met Trp
 20 25 30

<210> 58
 <211> 69
 <212> DNA
 <213> Sus scrofa

<400> 58
 tgggtacaagc acacggcgag tccccgctac cacacggtgg gccgcgccgc gggcctgctc 60
 atggggctg 69

<210> 59
 <211> 90
 <212> DNA
 <213> Sus scrofa

<400> 59
 tgggtacaagc acacggcgag tccccgctac cacacggtgg gccgcgccgc gggcctgctc 60
 atggggctgc gccgctcgcc ctacatgttg 90

<210> 60
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 60
 cgttctcggg gacataaacc ctg 23

<210> 61
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 61
 atgagcagcc cggaggcacg acc 23

<210> 62
 <211> 188
 <212> DNA
 <213> Rattus norvegicus

<400> 62

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```

ttcttgtcct aaccgcgcaa ggggccatgg acttgagcgc gctggcgctcg agcagagaag    60
tacggggccc tgggcccggg gctccgggtga accggcccct gctaccgcta ctgctgcttc    120
tgctcttgct acctctgccc gccagcgctt ggtacaagca cgtggcgagc cctcgctatc    180
acacagtg                                     188

```

<210> 63

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 63

atgagcagcc cggaggcacg acc 23

<210> 64

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 64

actgtgtgat agcgagggt cgc 23

<210> 65

<211> 615

<212> DNA

<213> Rattus norvegicus

<400> 65

```

ctcagagctg tactaggcag gaagagggac ggccctcagg gaaggggtggc cctatgctta    60
aaactttcct gtctcctctc cataagtgtc ccacttgtag caactcctac caagggggca    120
tcctttttgcc cctggcagcc catccttgta ttctgagacc atgcatggta ccagaactcc    180
ctccctgaca gttcccttcc tgggggagag gaaagggtaa gcaaggagat ccccccactaa    240
agcttcaagc gcagtccagc ttgcgatcta ctcatggga ggcttctagc taccggggtt    300
ccctcttctc cctccctctc catcctcctc tcccttgggc atgtgccgag ggggcgagcc    360
ggggcggggc cattgagaag ctgtagtcgc accaactgac tagtctcttc catcctccgg    420
agctccgacg ttctcgggga cataaaccct gttcttgtcc taaccgcgca agggggccatg    480
gacttgagcg cgctggcgtc gagcagagaa gtacggggcc ctgggcccgg ggctccgggtg    540
aaccggcccc tgctaccgct actgctgctt ctgctcttgc tacctctgcc cgccagcgcc    600
tggtacaagc acgtg                                     615

```

<210> 66

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 66

cgttctcggg gacataaacc ctg 23

<210> 67

<211> 24

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<212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 67
 cgagccctcg ctatcacaca gtgg 24

<210> 68
 <211> 497
 <212> DNA
 <213> Rattus norvegicus

<400> 68
 gtcgtgcctc cgggctgctc atggggctgc gccgctcgcc ctacctgtgg cgccgtgcct 60
 tgggtggggc cgctggaccg ctcgtggggc tcccgggaca gatggcccgc agcgctctcc 120
 tgcttccttc ccccgggcag gagctgtggg aggtacgaag caggagtcca ccggcaggac 180
 ttcccgctgca tgcaaccgag agtctgcggg acctggaggg agccggccaa cctgagcagt 240
 cgctaagctt tcagtcctgg acttcagcag agcccgtgc tagagccttc ggtgagacgc 300
 ttcgtgcccc gccatggttc ctgcagcaaa tcattcttgc cgatcctgtc aggctcgacg 360
 accgtctcaa gaaccgatgg cgccccgtg cttgacctaa gcaggagcac agcttgtagc 420
 tccagtcagg tctcgttgct tggccaataa aatcactctg attcccaaaa aaaaaaaaaa 480
 aaaaaaaaaa aaaaaaa 497

<210> 69
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 69
 ggggcggggc cattgagaag c 21

<210> 70
 <211> 21
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 70
 tgaccagaca acgagacctg a 21

<210> 71
 <211> 684
 <212> DNA
 <213> Rattus norvegicus

<400> 71
 tgtagtcgca ccaactgact agtctcttcc atcctccgga gctccgacgt tctcggggac 60
 ataaaccctg ttcttgtcct aaccgcgcaa ggggccatgg acttgagcgc gctggcgctc 120
 agcagagaag tacggggccc tggggccggg gctccggtga accggccccct gctaccgcta 180
 ctgctgcttc tgctcttgct acctctgccc gccagcgctt ggtacaagca cgtggcgagc 240
 cctcgctatc acacagtggg tcgtgcctcc gggctgctca tggggctgcg ccgctcgccc 300

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```

tacctgtggc gccgtgcctt ggggtggggcc gctggaccgc tcgtggggct cccgggacag 360
atggcccgc gcgtctctct gcttccttcc cccgggcagg agctgtggga ggtacgaagc 420
aggagtccac cggcaggact tcccgtgcat gcaaccgga gtctgcggga cctggaggga 480
gccggccaac ctgagcagtc gctaagcttt cagtcctgga cttcagcaga gcccgtgct 540
agagccttcg gtgagacgct tcgtgcccag ccatgggtcc tgcagcaaat catctttgcc 600
gatcctgtca ggctcgacga ccgtctcaag aaccgatggc gccccgtgc ttgacctaa 660
caggagcaca gctttagtct ccag 684

```

<210> 72

<211> 185

<212> PRT

<213> Rattus norvegicus

<400> 72

```

Met Asp Leu Ser Ala Leu Ala Ser Ser Arg Glu Val Arg Gly Pro Gly
1      5      10      15
Pro Gly Ala Pro Val Asn Arg Pro Leu Leu Pro Leu Leu Leu Leu
20     25     30
Leu Leu Leu Pro Leu Pro Ala Ser Ala Trp Tyr Lys His Val Ala Ser
35     40     45
Pro Arg Tyr His Thr Val Gly Arg Ala Ser Gly Leu Leu Met Gly Leu
50     55     60
Arg Arg Ser Pro Tyr Leu Trp Arg Arg Ala Leu Gly Gly Ala Ala Gly
65     70     75     80
Pro Leu Val Gly Leu Pro Gly Gln Met Ala Arg Ser Ala Leu Leu Leu
85     90     95
Pro Ser Pro Gly Gln Glu Leu Trp Glu Val Arg Ser Arg Ser Ser Pro
100    105    110
Ala Gly Leu Pro Val His Ala Thr Arg Ser Leu Arg Asp Leu Glu Gly
115    120    125
Ala Gly Gln Pro Glu Gln Ser Leu Ser Phe Gln Ser Trp Thr Ser Ala
130    135    140
Glu Pro Ala Ala Arg Ala Phe Gly Glu Thr Leu Arg Ala Gln Pro Trp
145    150    155    160
Phe Leu Gln Gln Ile Phe Ala Asp Pro Val Arg Leu Asp Asp Arg
165    170    175
Leu Lys Asn Arg Trp Arg Pro Arg Ala
180    185

```

<210> 73

<211> 23

<212> PRT

<213> Rattus norvegicus

<400> 73

```

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1      5      10      15
Ser Gly Leu Leu Met Gly Leu
20

```

<210> 74

<211> 30

<212> PRT

<213> Rattus norvegicus

<400> 74

```

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1      5      10      15
Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Leu Trp

```

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20

25

30

<210> 75
 <211> 69
 <212> DNA
 <213> Rattus norvegicus

<400> 75
 tggtagaagc acgtggcgag ccctcgctat cacacagtgg gtcgtgcctc cgggctgctc 60
 atggggctg 69

<210> 76
 <211> 90
 <212> DNA
 <213> Rattus norvegicus

<400> 76
 tggtagaagc acgtggcgag ccctcgctat cacacagtgg gtcgtgcctc cgggctgctc 60
 atggggctgc gccgctcgcc ctacctgtgg 90

<210> 77
 <211> 23
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Probe

<400> 77
 ttcacacctca acctggccat cgc 23

<210> 78
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 78
 acccagttct tgcctaacc ctcc 24

<210> 79
 <211> 24
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer

<400> 79
 cctgcttcgt acctcccaca gctc 24

<210> 80
 <211> 311
 <212> DNA
 <213> Mus musculus

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<400> 80
aaggggcaat tgacgtgagc gcgctggcgt ctaacagaga agtacggggc cctgggcccc 60
ggactcccag gaaccggccc ctgctgcccc tgctgctgct tctgctcttg ctaccgctgc 120
ccgccagcgc ctggtataag cacgtggcga gtccccgcta tcacacagtg ggtcgtgcct 180
ccgggctgct catggggctg cgccgctcgc cctaccagtg gcgccgtgcc ctgggcgggg 240
ctgctggacc cctctcccgg ctcccaggac cggtcgcccc cggcgctctc ctgcttcctt 300
cctcagggca g 311

<210> 81
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 81
catgagcagc ccggaggcac gacc 24

<210> 82
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 82
gtgatagcgg ggactcgcca cgtg 24

<210> 83
<211> 237
<212> DNA
<213> Mus musculus

<400> 83
aaaggctgta gtgcaccaa ctgactggtc tccatcctct ggagctccga cgtgctcggt 60
ctcggagaca taaaccagc tcttgctcta accctccaag gggcaattga cgtgagcgcg 120
ctggcgtcta acagagaagt acggggccct gggcccgga ctcccaggaa ccggcccctg 180
ctgcccctgc tgctgcttct gctcttgcta ccgctgcccc ccagcgcttg gtataag 237

<210> 84
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 84
accagttct tgtcctaacc ctcc 24

<210> 85
<211> 24
<212> DNA
<213> Artificial Sequence

<220>

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<223> Primer

<400> 85

gggcaattga cgtgagcgcg ctgg 24

<210> 86

<211> 598

<212> DNA

<213> Mus musculus

<400> 86

cgtctaacag	agaagtacgg	ggccctgggc	ccgggactcc	caggaaccgg	cccctgctgc	60
ccctgctgct	gcttctgctc	ttgctaccgc	tgcccgccag	cgcttggtat	aagcacgtgg	120
cgagtcctccg	ctatcacaca	gtgggtcgtg	cctccgggct	gctcatgggg	ctgcgccgct	180
cgccctacca	gtggcgccgt	gccctgggcg	gggctgctgg	acccctctcc	cggctcccag	240
gaccggctcg	ccgcggcgct	ctcctgcttc	cttcctcagg	gcaggagctg	tgggaggtac	300
gaagcaggag	ctcacctgca	gggcttcccg	tccatgcacc	ctggagtccg	cgggacctgg	360
agggagtccg	ccaaccggag	cagtcgctaa	gccttcactc	ctggatctca	gaggagcccg	420
ctgctagagc	cttcggagag	acgcttcgtg	cccagccatg	gttcctgcag	caagtcattc	480
ttgccgatcc	tgtcaggccc	aagaaccgat	ggcgcccca	tgcttgacct	aggcaggagc	540
acagcttgaa	gctccagtca	ggcctcgtgt	ttctgggtcaa	taaaaccaac	ctgattcc	598

<210> 87

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 87

aaaggctgta gtcgcaccaa c 21

<210> 88

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 88

accagaaaca cgaggcctga c 21

<210> 89

<211> 659

<212> DNA

<213> Mus musculus

<400> 89

tgactggtct	ccatcctctg	gagctccgac	gtgctcgttc	tggagacat	aaaccagtt	60
cttgtcctaa	ccctccaagg	ggcaattgac	gtgagcgcg	tggcgtctaa	cagagaagta	120
cggggccctg	ggcccgggac	tcccaggaac	cggcccctgc	tggccctgct	gctgcttctg	180
ctcttgctac	cgctgcccg	cagcgccctg	tataagcacg	tggcgagtcc	ccgctatcac	240
acagtgggtc	gtgcctccgg	gctgctcatg	gggctgcgcc	gctcgcccta	ccagtggcgc	300
cgtgccctgg	gcggggctgc	tggacccctc	tcccggctcc	caggaccggg	cgcccgcggc	360
gctctcctgc	ttccttcttc	agggcaggag	ctgtgggagg	tacgaagcag	gagctcacct	420
gcagggtctc	ccgtccatgc	accctggagt	ccgcgggacc	tggagggagt	ccgccaaccg	480

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```

gagcagtcgc taagccttca ctccctggatc tcagaggagc ccgctgctag agccttcgga 540
gagacgcttc gtgcccagcc atggttcctg cagcaagtca tctttgccga tcctgtcagg 600
cccaagaacc gatggcgccc ccatgcttga cctaggcagg agcacagctt gaagctcca 659

```

<210> 90
 <211> 176
 <212> PRT
 <213> Mus musculus

```

<400> 90
Leu Ala Ser Asn Arg Glu Val Arg Gly Pro Gly Pro Gly Thr Pro Arg
1          5          10          15
Asn Arg Pro Leu Leu Pro Leu Leu Leu Leu Leu Leu Leu Pro Leu
20          25          30
Pro Ala Ser Ala Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr
35          40          45
Val Gly Arg Ala Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr
50          55          60
Gln Trp Arg Arg Ala Leu Gly Gly Ala Ala Gly Pro Leu Ser Arg Leu
65          70          75          80
Pro Gly Pro Val Ala Arg Gly Ala Leu Leu Leu Pro Ser Ser Gly Gln
85          90          95
Glu Leu Trp Glu Val Arg Ser Arg Ser Ser Pro Ala Gly Leu Pro Val
100         105         110
His Ala Pro Trp Ser Pro Arg Asp Leu Glu Gly Val Arg Gln Pro Glu
115         120         125
Gln Ser Leu Ser Leu His Ser Trp Ile Ser Glu Glu Pro Ala Ala Arg
130         135         140
Ala Phe Gly Glu Thr Leu Arg Ala Gln Pro Trp Phe Leu Gln Gln Val
145         150         155         160
Ile Phe Ala Asp Pro Val Arg Pro Lys Asn Arg Trp Arg Pro His Ala
165         170         175

```

<210> 91
 <211> 23
 <212> PRT
 <213> Mus musculus

```

<400> 91
Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1          5          10          15
Ser Gly Leu Leu Met Gly Leu
20

```

<210> 92
 <211> 30
 <212> PRT
 <213> Mus musculus

```

<400> 92
Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1          5          10          15
Ser Gly Leu Leu Met Gly Leu Arg Arg Ser Pro Tyr Gln Trp
20          25          30

```

<210> 93
 <211> 69
 <212> DNA

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<213> Mus musculus

<400> 93

tggtataagc acgtggcgag tccccgctat cacacagtgg gtcgtgcctc cgggctgctc 60
atggggctg 69

<210> 94

<211> 90

<212> DNA

<213> Mus musculus,

<400> 94

tggtataagc acgtggcgag tccccgctat cacacagtgg gtcgtgcctc cgggctgctc 60
atggggctgc gccgctcgcc ctaccagtgg 90

<210> 95

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> Human GPR8 ligand (1-23) oxidant

<222> 21

<223> Xaa is Met(O)

<400> 95

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1 5 10 15
Ala Gly Leu Leu Xaa Gly Leu
20

<210> 96

<211> 22

<212> PRT

<213> Homo sapiens

<400> 96

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1 5 10 15
Ala Gly Leu Leu Met Gly
20

<210> 97

<211> 21

<212> PRT

<213> Homo sapiens

<400> 97

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1 5 10 15
Ala Gly Leu Leu Met
20

<210> 98

<211> 20

<212> PRT

<213> Homo sapiens

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<400> 98

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala Gly Leu Leu
 20

<210> 99

<211> 19

<212> PRT

<213> Homo sapiens

<400> 99

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala Gly Leu

<210> 100

<211> 18

<212> PRT

<213> Homo sapiens

<400> 100

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala Gly

<210> 101

<211> 17

<212> PRT

<213> Homo sapiens

<400> 101

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala

<210> 102

<211> 16

<212> PRT

<213> Homo sapiens

<400> 102

Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15

<210> 103

<211> 23

<212> PRT

<213> Artificial Sequence

<220>

<221> Porcine GPR8 ligand (1-23) oxidant

<222> 21

<223> Xaa is Met(O)

<400> 103

Trp Tyr Lys His Thr Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15

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Ala Gly Leu Leu Xaa Gly Leu
20

<210> 104
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<221> Rat/mouse GPR8 ligand (1-23) oxidant
<222> 21
<223> Xaa is Met(O)

<400> 104
Trp Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1 5 10 15
Ser Gly Leu Leu Xaa Gly Leu
20

<210> 105
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<221> Fmoc-added human GPR8L (1-23)
<222> 1
<223> Xaa is Fmoc-Trp

<400> 105
Xaa Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1 5 10 15
Ala Gly Leu Leu Met Gly Leu
20

<210> 106
<211> 23
<212> PRT
<213> Artificial Sequence

<220>
<221> [N α -Acetyl-Trp1]-human GPR8 ligand (1-23)
<222> 1
<223> Xaa is Ac-Trp

<400> 106
Xaa Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
1 5 10 15
Ala Gly Leu Leu Met Gly Leu
20

<210> 107
<211> 22
<212> PRT
<213> Homo sapiens

<400> 107
Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala Ala

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1 5 10 15
 Gly Leu Leu Met Gly Leu
 20

<210> 108
 <211> 20
 <212> PRT
 <213> Homo sapiens

<400> 108
 His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala Ala Gly Leu
 1 5 10 15
 Leu Met Gly Leu
 20

<210> 109
 <211> 15
 <212> PRT
 <213> Homo sapiens

<400> 109
 Arg Tyr His Thr Val Gly Arg Ala Ala Gly Leu Leu Met Gly Leu
 1 5 10 15

<210> 110
 <211> 9
 <212> PRT
 <213> Homo sapiens

<400> 110
 Arg Ala Ala Gly Leu Leu Met Gly Leu
 1 5

<210> 111
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> [N-Acetyl-Tyr2]-human GPR8 ligand (2-23)
 <222> 1
 <223> Xaa is Ac-Tyr

<400> 111
 Xaa Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala Ala
 1 5 10 15
 Gly Leu Leu Met Gly Leu
 20

<210> 112
 <211> 23
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> [D-Trp1]-human GPR8 ligand (1-23)
 <222> 1
 <223> Xaa is D-Trp

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<400> 112
 Xaa Tyr Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala
 1 5 10 15
 Ala Gly Leu Leu Met Gly Leu
 20

<210> 113
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <221> [N-3-Indolepropanoyl-Tyr2]-human GPR8 ligand (2-23)
 <222> 1
 <223> Xaa is 3-Indolepropanoyl-Tyr

<400> 113
 Xaa Lys His Val Ala Ser Pro Arg Tyr His Thr Val Gly Arg Ala Ala
 1 5 10 15
 Gly Leu Leu Met Gly Leu
 20

<210> 114
 <211> 66
 <212> DNA
 <213> Homo sapiens

<400> 114
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 atgggg 66

<210> 115
 <211> 63
 <212> DNA
 <213> Homo sapiens

<400> 115
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 atg 63

<210> 116
 <211> 60
 <212> DNA
 <213> Homo sapiens

<400> 116
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<210> 117
 <211> 57
 <212> DNA
 <213> Homo sapiens

<400> 117
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<210> 118

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<211> 54
 <212> DNA
 <213> Homo sapiens

<400> 118
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<210> 119
 <211> 51
 <212> DNA
 <213> Homo sapiens

<400> 119
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<210> 120
 <211> 48
 <212> DNA
 <213> Homo sapiens

<400> 120
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<210> 121
 <211> 66
 <212> DNA
 <213> Homo sapiens

<400> 121
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 gggctg 66

<210> 122
 <211> 60
 <212> DNA
 <213> Homo sapiens

<400> 122
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<210> 123
 <211> 45
 <212> DNA
 <213> Homo sapiens

<400> 123
 cgctaccaca cggtggggcg cgccgctggc ctgctcatgg ggctg 45

<210> 124
 <211> 27
 <212> DNA
 <213> Homo sapiens

<400> 124
 cgccgccgctg gcctgctcat ggggctg 27

<210> 125
 <211> 51

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<212> DNA
 <213> *Sus scrofa*

<400> 125
 tgggtacaagc acacggcgag tccccgctac cacacggtgg gccgcgccgc g

51

<210> 126
 <211> 329
 <212> PRT
 <213> *Rattus norvegicus*

<400> 126
 Met His Asn Leu Ser Leu Phe Glu Pro Gly Arg Gly Asn Val Ser Cys
 5 10 15
 Gly Gly Pro Phe Leu Gly Cys Pro Asn Glu Ser Asn Pro Ala Pro Leu
 20 25 30
 Pro Leu Pro Gln Pro Leu Ala Val Ala Val Pro Val Val Tyr Gly Val
 35 40 45
 Ile Cys Ala Val Gly Leu Ala Gly Asn Ser Ala Val Leu Tyr Val Leu
 50 55 60
 Leu Arg Thr Pro Arg Met Lys Thr Val Thr Asn Val Phe Ile Leu Asn
 65 70 75 80
 Leu Ala Ile Ala Asp Glu Leu Phe Thr Leu Val Leu Pro Ile Asn Ile
 85 90 95
 Ala Asp Phe Leu Leu Arg Arg Trp Pro Phe Gly Glu Val Met Cys Lys
 100 105 110
 Leu Ile Val Ala Val Asp Gln Tyr Asn Thr Phe Ser Ser Leu Tyr Phe
 115 120 125
 Leu Ala Val Met Ser Ala Asp Arg Tyr Leu Val Val Leu Ala Thr Ala
 130 135 140
 Glu Ser Arg Arg Val Ser Gly Arg Thr Tyr Gly Ala Ala Arg Ala Val
 145 150 155 160
 Ser Leu Ala Val Trp Ala Leu Val Thr Leu Val Val Leu Pro Phe Ala
 165 170 175
 Val Phe Ala Arg Leu Asp Glu Glu Gln Gly Arg Arg Gln Cys Val Leu
 180 185 190
 Val Phe Pro Gln Pro Glu Ala Phe Trp Trp Arg Ala Ser Arg Leu Tyr
 195 200 205
 Thr Leu Val Leu Gly Phe Ala Ile Pro Val Ser Thr Ile Cys Ala Leu
 210 215 220
 Tyr Ile Thr Leu Leu Cys Arg Leu Arg Ala Ile Gln Leu Asp Ser His
 225 230 235 240
 Ala Lys Ala Leu Asp Arg Ala Lys Lys Arg Val Thr Leu Leu Val Val
 245 250 255
 Ala Ile Leu Ala Val Cys Leu Leu Cys Trp Thr Pro Tyr His Leu Ser
 260 265 270
 Thr Ile Val Ala Leu Thr Thr Asp Leu Pro Gln Thr Pro Leu Val Ile
 275 280 285
 Gly Ile Ser Tyr Phe Ile Thr Ser Leu Ser Tyr Ala Asn Ser Cys Leu
 290 295 300
 Asn Pro Phe Leu Tyr Ala Phe Leu Asp Asp Ser Phe Arg Arg Ser Leu
 305 310 315 320
 Arg Gln Leu Val Ser Cys Arg Thr Ala
 325

<210> 127
 <211> 987
 <212> DNA

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<213> *Rattus norvegicus*

<400> 127

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atgcacaact tgtcgctctt cgagcctggc aggggcaatg tgtcttgcg cgccccattt 60
ttgggctgtc ctaacgagtc gaacccagcg cctctgccac tgccgcagcc tctggcggtg 120
gcagtgcctg tgggtctacgg ggtgatctgc gcgggtggac tggcgggcaa ctccgcggtg 180
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agtctggcgg tgtggcgct ggtgacattg gtgctgctgc cttttgcggt attcgcccgg 540
ctggacgaag agcagggtcg gcgtcagtc gtgctggtct tcccgagcc tgaggccttc 600
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<210> 128

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 128

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actgatatgc acaacttgct gctcttcg 28

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<210> 129

<211> 28

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 129

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actagtccag gctgtgcggc atgacacc 28

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<210> 130

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 130

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gttggtggtg gcgattctg 19

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<210> 131

<211> 19

<212> DNA

<213> Artificial Sequence

- 31 / 36 -

<220>

<223> Primer

<400> 131

tggtgagcgc cactatggt 19

<210> 132

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 132

gtccgcgatg ttgatgggca gcac 24

<210> 133

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 133

gaagagctca tcggcgatag ccag 24

<210> 134

<211> 440

<212> DNA

<213> Mus musculus

<400> 134

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cgcggggctg	ctgagtggaa	tcctgggtgg	cgctgtctct	ccagccctct	ccaagatgca	180
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ctgtcccaac	gggtccagcc	tggtctctct	gccgctgccg	cagccactgg	cggtagcagt	300
gcctgtcgct	tacggggtaa	tttgcgccgt	gggactggct	ggcaactctg	cggtgctgta	360
cgtactgctg	cgcacgccgc	gcatgaagac	tgtcaccaac	gtgttcaccc	tcaacctggc	420
tatcgccgat	gagctcttca					440

<210> 135

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 135

tttcgcgggg ctgctgagtg gaat 24

<210> 136

<211> 24

<212> DNA

- 32 / 36 -

<213> Artificial Sequence

<220>

<223> Primer

<400> 136

agtgtctgcct gcggtggaaa gagg 24

<210> 137

<211> 1083

<212> DNA

<213> Mus musculus

<400> 137

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gcataactta acgcttttcg agtctggagg ggacaacgtg tcttgccggc gctcatcttt 120
gggctgtccc aacgggtcca gcctggctcc tctgcccgtg ccgcagccac tggcggtagc 180
agtgcctgtc gtctacgggg taatttgccg cgtgggactg gctggcaact ctgcggtgct 240
gtacgtactg ctgcgcacgc cgcgcacatgaa gactgtcacc aacgtgttca tcctcaacct 300
ggctatcgcc gatgagctct tcacctcgt gctgcccata aacatcgagg acttcctgct 360
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cactttctct agcctctact tcctcgccgt catgagcgcc gaccgatacc tggtggttct 480
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gtggcgtgcc agccgtctct acacactagt attgggcttt gccatcccgg tgaccacat 720
ctgtgctctc tataccactc tgctctgccc actgcgtgct atccagctag atagccacgc 780
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cccgcaaacg ccgctgggtc tcggcatctc ttacttcata accagcctga gctatgctaa 960
cagctgcctc aaccctttcc tctatgcctt cctggacgac agcttccgca gaagcctccg 1020
gcaattggtg tcatgccgtt cagcctgatg ccctttccac ctctttccac cgcaggcagc 1080
act 1083

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<210> 138

<211> 329

<212> PRT

<213> Mus musculus

<400> 138

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Met His Asn Leu Thr Leu Phe Glu Ser Gly Gly Asp Asn Val Ser Cys
      5              10              15
Gly Gly Ser Ser Leu Gly Cys Pro Asn Gly Ser Ser Leu Ala Pro Leu
      20              25              30
Pro Leu Pro Gln Pro Leu Ala Val Ala Val Pro Val Val Tyr Gly Val
      35              40              45
Ile Cys Ala Val Gly Leu Ala Gly Asn Ser Ala Val Leu Tyr Val Leu
      50              55              60
Leu Arg Thr Pro Arg Met Lys Thr Val Thr Asn Val Phe Ile Leu Asn
      65              70              75              80
Leu Ala Ile Ala Asp Glu Leu Phe Thr Leu Val Leu Pro Ile Asn Ile
      85              90              95
Ala Asp Phe Leu Leu Arg Arg Trp Pro Phe Gly Glu Val Met Cys Lys
      100             105             110
Leu Ile Val Ala Val Asp Gln Tyr Asn Thr Phe Ser Ser Leu Tyr Phe
      115             120             125
Leu Ala Val Met Ser Ala Asp Arg Tyr Leu Val Val Leu Ala Thr Ala
      130             135             140

```


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Glu Ser Arg Arg Val Ser Gly Arg Thr Tyr Gly Ala Ala Arg Ala Val
 145 150 155 160
 Ser Leu Ala Val Trp Ala Leu Val Thr Leu Val Val Leu Pro Phe Ala
 165 170 175
 Val Phe Ala Arg Leu Asp Glu Glu Gln Gly Arg Arg Gln Cys Val Leu
 180 185 190
 Val Phe Pro Gln Pro Glu Ala Phe Trp Trp Arg Ala Ser Arg Leu Tyr
 195 200 205
 Thr Leu Val Leu Gly Phe Ala Ile Pro Val Thr Thr Ile Cys Ala Leu
 210 215 220
 Tyr Thr Thr Leu Leu Cys Arg Leu Arg Ala Ile Gln Leu Asp Ser His
 225 230 235 240
 Ala Lys Ala Leu Asp Arg Ala Lys Lys Arg Val Thr Leu Leu Val Ala
 245 250 255
 Ala Ile Leu Ala Val Cys Leu Leu Cys Trp Thr Pro Tyr His Leu Ser
 260 265 270
 Thr Ile Val Ala Leu Thr Thr Asp Leu Pro Gln Thr Pro Leu Val Ile
 275 280 285
 Gly Ile Ser Tyr Phe Ile Thr Ser Leu Ser Tyr Ala Asn Ser Cys Leu
 290 295 300
 Asn Pro Phe Leu Tyr Ala Phe Leu Asp Asp Ser Phe Arg Arg Ser Leu
 305 310 315 320
 Arg Gln Leu Val Ser Cys Arg Ser Ala
 325

<210> 139

<211> 987

<212> DNA

<213> Mus musculus

<400> 139

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gcagtgcctg	tcgtctacgg	ggtaatttgc	gccgtgggac	tggttgga	ctctgcgggtg	180
ctgtacgtac	tgctgcgcac	gccgcgcacg	aagactgtca	ccaacgtgtt	catcctcaac	240
ctggctatcg	ccgatgagct	cttcaccctc	gtgctgcca	tcaacatcgc	ggacttcctg	300
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aacactttct	ctagcctcta	cttcctcgcc	gtcatgagcg	cggaccgata	cctgggtggtt	420
ctggccacag	cagagtcgcg	ccgggtgtcc	gggcgcactt	acgggtgcagc	gcgtgctgtc	480
agtctggcgg	tgtgggcgct	ggtgacgctg	gtcgtgctgc	cctttgcggg	attcgctcgg	540
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gccaaaggccc	tggtatcgtgc	caagaagcgc	gtgaccttgt	tggtggcggc	gattctggct	780
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ctcccga	cgccgtgggt	catcggcac	tcttacttca	tcaccagcct	gagctatgct	900
aacagctgcc	tcaacccttt	cctctatgcc	ttcctggacg	acagcttcg	cagaagcctc	960
cggcaattgg	tgtcatgccg	ttcagcc				987

<210> 140

<211> 27

<212> DNA

<213> Artificial Sequence

<220>

<223> Probe

<400> 140

- 34 / 36 -

tcctctgctg gacaccgtac cacctga 27

<210> 141

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 141

atcgatatgg acaacgcctc gttctcggag cc 32

<210> 142

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 142

actagtgtca ggctgccgcg cggcaagtta tc 32

<210> 143

<211> 1000

<212> DNA

<213> Homo sapiens

<400> 143

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atcgatatgg acaacgcctc gttctcggag ccctggcccc ccaacgcata gggcccggac 60
ccggcgctga gctgctccaa cgcgtcgact ctggcgccgc tgccggcgcc gctggcggtg 120
gctgtaccag ttgtctacgc ggtgatctgc gccgtgggtc tggcgggcaa ctccgccgtg 180
ctgtacgtgt tgetgcgggc gccccgcata aagaccgtca ccaacctgtt catcctcaac 240
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aacagctgcc tcaaccctt cctctacgcc ttcttgagcg ccagcttccg caggaaacctc 960
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<210> 144

<211> 328

<212> PRT

<213> Homo sapiens

<400> 144

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Met Asp Asn Ala Ser Phe Ser Glu Pro Trp Pro Ala Asn Ala Ser Gly
1           5           10          15
Pro Asp Pro Ala Leu Ser Cys Ser Asn Ala Ser Thr Leu Ala Pro Leu
          20          25          30

```

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Pro Ala Pro Leu Ala Val Ala Val Pro Val Val Tyr Ala Val Ile Cys
    35              40              45
Ala Val Gly Leu Ala Gly Asn Ser Ala Val Leu Tyr Val Leu Leu Arg
    50              55              60
Ala Pro Arg Met Lys Thr Val Thr Asn Leu Phe Ile Leu Asn Leu Ala
65              70              75              80
Ile Ala Asp Glu Leu Phe Thr Leu Val Leu Pro Ile Asn Ile Ala Asp
    85              90              95
Phe Leu Leu Arg Gln Trp Pro Phe Gly Glu Leu Met Cys Lys Leu Ile
    100             105             110
Val Ala Ile Asp Gln Tyr Asn Thr Phe Ser Ser Leu Tyr Phe Leu Thr
    115             120             125
Val Met Ser Ala Asp Arg Tyr Leu Val Val Leu Ala Thr Ala Glu Ser
    130             135             140
Arg Arg Val Ala Gly Arg Thr Tyr Ser Ala Ala Arg Ala Val Ser Leu
145             150             155             160
Ala Val Trp Gly Ile Val Thr Leu Val Val Leu Pro Phe Ala Val Phe
    165             170             175
Ala Arg Leu Asp Asp Glu Gln Gly Arg Arg Gln Cys Val Leu Val Phe
    180             185             190
Pro Gln Pro Glu Ala Phe Trp Trp Arg Ala Ser Arg Leu Tyr Thr Leu
    195             200             205
Val Leu Gly Phe Ala Ile Pro Val Ser Thr Ile Cys Val Leu Tyr Thr
    210             215             220
Thr Leu Leu Cys Arg Leu His Ala Met Arg Leu Asp Ser His Ala Lys
225             230             235             240
Ala Leu Glu Arg Ala Lys Lys Arg Val Thr Phe Leu Val Val Ala Ile
    245             250             255
Leu Ala Val Cys Leu Leu Cys Trp Thr Pro Tyr His Leu Ser Thr Val
    260             265             270
Val Ala Leu Thr Thr Asp Leu Pro Gln Thr Pro Leu Val Ile Ala Ile
    275             280             285
Ser Tyr Phe Ile Thr Ser Leu Ser Tyr Ala Asn Ser Cys Leu Asn Pro
    290             295             300
Phe Leu Tyr Ala Phe Leu Asp Ala Ser Phe Arg Arg Asn Leu Arg Gln
305             310             315             320
Leu Ile Thr Cys Arg Ala Ala Ala
    325

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<210> 145

<211> 32

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 145

atcgatatgg acaacgcctc gttctcggag cc 32

<210> 146

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

- 36 / 36 -

<400> 146
tagaggctgg agaaggtggt g 21

<210> 147
<211> 21
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 147
catgaagacc gtcaccaacc t 21

<210> 148
<211> 19
<212> DNA
<213> Artificial Sequence

<220>
<223> Primer

<400> 148
ccagcgtgaa gagctcgtc 19

<210> 149
<211> 20
<212> PRT
<213> Artificial Sequence

<220>
<223> [Phe2] human GPR8 ligand (1-20)

<400> 149
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Ala Gly Leu Leu
20

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<213> Artificial Sequence

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<223> Sequence encoding [Phe2] human GPR8 ligand (1-20)

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